February 18, 2004, 14:12:09;

); Search time 6.5921 Seconds (without alignments) 87.531 Million cell updates/sec

Title: Perfect score: Sequence: US-09-643-260-3 26 LDASAL 6

Scoring table: BLOSUM62 Gapop 10.0 , Gapext 0.5

Searched: 283308 segs, 96168682 residues

Total number of hits satisfying chosen parameters:

Minimum DB Maximum DB length: 0 length: 2000000000

Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries

PIR_76:* 1: pir1:* pir2:* pir3:* pir4:*

and is No. is the number of results predicted by chance to have a greater than or equal to the score of the result being printed, derived by analysis of the total score distribution.

SUMMARIES

| 29 | 28 | 27 | 26 | 25 | 24 | 23 | 22 | 21 | 20 | 19 | 18 | 17 | 16 | 15 | 14 | 13 | 12 | 11 | 10 | 9 | 00 | 7 | σ | ທ | 4. | ω | N | بر | Result No. |
|-------------------|--------------------|--------------------|--------------------|--------------------|--------------------|--------------------|--------|-------------------|--------|---------|--------------------|--------------------|--------------------|--------------------|-----|--------|--------------------|--------|--------|--------|--------------------|-----|--------------------|---------------|--------------------|-------|--------|--------|---------------|
| 24 | 24 | 24 | 24 | 26 | 26 | 26 | 26 | 26 | 26 | 26 | 26 | 26 | 26 | 26 | 26 | 26 | 26 | 26 | 26 | 26 | 26 | 26 | 26 | 26 | 26 | 26 | 26 | 26 | Score |
| | 92.3 | 'n | | .0 | | .0 | | | .0 | | | 100.0 | | | | | | | | | | | | | | 100.0 | | 100.0 | Query |
| 179 | 166 | 157 | 58 | 1313 | 1006 | 920 | 894 | 855 | 638 | 586 | 550 | 516 | 513 | 513 | 512 | 483 | 437 | 394 | 394 | 383 | 334 | 281 | 259 | 230 | 171 | 130 | 129 | 841 | Length |
| N | N | N | ` | | | | | | | | | N | | | | | | | | | | | | | | | | ٠., | B |
| B96989 | AC1940 | C70882 | 855558 | JC2038 | T41439 | 140614 | G82260 | T41336 | 139196 | T49210 | H70772 | E81092 | AH3019 | A96265 | 84 | AH3265 | A70587 | B81062 | H81807 | H98287 | T37024 | | F69311 | E95326 | 8762 | 10 | 3120 | D70672 | ID |
| probable membrane | purine-binding che | hypothetical prote | allophycocyanin be | peptidyl-dipeptida | putitive sulfite r | surface array prot | 댞 | probable nitrogen | e sei | cal pro | probable argS prot | hypothetical prote | sigma 54 dependent | hypothetical prote | Ω | 3 | hypothetical prote | | | ical | probable DNA-bindi | cal | conserved hypothet | | hypothetical prote | ጟ | ical [| o | Description |

| 4 5 | 44 | 43 | 42 | 41 | 40 | 39 | 38 | 37 | 36 | 35 | 34 | 33 | 32 | 31 | 30 | |
|--------------------|--------------------|--------------------|--------------------|--------------------|--------------------|--------------------|--------------------|--------------------|--------------------|--------------------|--------------------|--------------------|--------------------|--------------------|--------------------|--|
| 24 | 24 | 24 | 24 | 24 | 24 | 24 | 24 | 24 | 24 | 24 | 24 | 24 | 24 | 24 | 24 | |
| 92.3 | 92.3 | 92.3 | 92.3 | 92.3 | 92.3 | 92.3 | 92.3 | 92.3 | 92.3 | 92.3 | 92.3 | 92.3 | 92.3 | 92.3 | 92.3 | |
| 405 | 401 | 393 | 384 | 359 | 359 | 359 | 346 | 326 | 304 | 298 | 294 | 292 | 292 | 279 | 197 | |
| ۲ | N | N | N | N | N | Н | Н | N | N | N | N | N | N | N | N | |
| XUECSD | AC2113 | 869191 | G85948 | AC0875 | D91103 | ADEC2A | I78840 | T09995 | T42939 | A41227 | T26946 | H98028 | A95163 | A83986 | A64484 | |
| dihydrolipoamide S | alanine racemase [| adenosylmethionine | fructose-bisphosph | fructose 1,6-bisph | fructose-bisphosph | fructose-bisphosph | protein kinase (EC | phosphoprotein pho | hypothetical prote | protein kinase (EC | hypothetical prote | hypothetical prote | hypothetical prote | hypothetical prote | conserved hypothet | |

ALIGNMENTS

hypothetical protein Rv2975c - Mycobacterium tuberculosis (strain H37RV)
C;Species: Mycobacterium tuberculosis
C;Date: 17-Jull-1998 #sequence_revision 17-Jul-1998 #text_change 22-Oct-1999
C;Accession: D70672
R;Accession: D70672
R;Cole, S.T.; Brosch, R.; Parkhill, J.; Garnier, T.; Churcher, C.; Harris, D.; Gord; Connor, R.; Davies, R.; Devlin, K.; Feltwell, T.; Gentles, S.; Hamlin, N.; Holroy Rajandream, M.A.; Rogers, J.; Rutter, S.; Seeger, K.; Skelton, S.; Squares, S.
Nature 393, 537-544, 1998 A;Authors: Sqares, R.; Sulston, J.E.; Taylor, K.; Whitehead, S.; Barrell A;Title: Deciphering the biology of Mycobacterium tuberculosis from the A;Reference number: A70500; MUID:98295987; PMID:9634230 A;Accession: D70672 A;Cross-references: GB:Z83018; GB:AL123456; NID:g3261671; PIDN:CAB05437.1; PID:e283 A;Experimental source: strain H37Rv A; Molecule type: DNA A; Residues: 1-84 <COL> A;Gene: Rv2975c A; Status: preliminary; nucleic acid sequence not shown; translation not Query Match
Best Local Similarity
Matches 6; Conserv Genetics: 100.0%; Score 26; DB 100.0%; Pred. No. 12; Live 0; Mismatches 2; Length S.; Barrell, complete B.G.

ge

밁 Ş ω 1 LDASAL 6 LDASAL 13

Conservative

0

Indels

0;

Gaps

0

A; Molecule type: \A; Residues: 1-129 A;Status: preliminary; translated from GB/EMBL/DDBJA;Molecule type: VNA A; Accession: A; Description hetical netical protein 633 - Sphingomonas aromaticivorans plasmid pNL1
sies: Sphingomonas aromaticivorans
11-Jan-2000 #sequence_revision 11-Jan-2000 #text_change 11-Jan-2000 **3**.77. F; Stillwell, L.C.; Wong, K.K.; Thurston, S.J.; Sisk, E.C.; Sensen, C the EMBL Data Library, July 1998 c. Complete sequence of a 184 kb catabolic plasmid from Sphingomonas a Number: Z20992 ROM> \EMBL:AF079317; NID:g3378261; PID:g3378341; PIDN:AAD03924.1

A; Genome: plasmid pNL A; Note: orf633

Genetics: Cross-references:

Query Match Best Local Similarity 100.0%; Score 26; DB 2; Pred. No. 19; Length 129;

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Minimum
Maximum
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Perfect score:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Post-processing:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Searched:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DB seq
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length: 2000000000
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40
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         A_Geneseq_19Jun03:*
1: /SIDS1/qcqdat=/~
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         BLOSUM62
Gapop 10.0 , Gapext 0.5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               February 18, 2004, 13:37:19;
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                     Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
| SIDS1/gcgdata/geneseq/geneseqp-embl/AA1980.DAT:*
| SIDS1/gcgdata/geneseq/geneseqp-embl/AA1981.DAT:*
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Copyright
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GenCore version 5.1.6 (c) 1993 - 2004 Compugen Ltd.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ; Search time 22.7763 Seconds (without alignments) 41.814 Million cell updates/sec
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

| 9 | 8 | 7 | 9 | (J) | 4 | ω | 2 | ь | Result No. |
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| 40 | 40 | 40 | 40 | 40 | 40 | 40 | 40 | 40 | Score |
| 100.0 | 100.0 | 100.0 | 100.0 | 100.0 | 100.0 | 100.0 | 100.0 | 100.0 | % Query Match |
| ø | 9 | 80 | œ | 7 | 0 | o, | o, | οn | % Query Match Length DB |
| 23 | 20 | 23 | 23 | 23 | 24 | 23 | 23 | 23 | BB |
| AAM48526 | AAW96182 ∀_ | AAM48535 | AAM48527 | AAM48534 | ABU08418 | AAM48655 | AAM48530 | ABB08725 | ij |
| Anti-inflammatory | IKK-alpha polypept | Anti-inflammatory | Anti-inflammatory | Anti-inflammatory | Human NEMO binding | NBD mutant peptide | Anti-inflammatory | IKKbeta NEMO bindi | Description |

| 45 | 44 | 43 | 42 | 41 | 40 | 39 | 38 | 37 | 3 6 | ა წ | 34 | ω ω | 32 | 31 | 30 | 29 | 28 | 27 | 26 | 25 | 24 | 23 | 22 | 21 | 20 | 19 | 18 | 17 | 16 | 15 | 14 | 13 | 12 | 11 | 10 |
|--------------------|----------|----------|--------------------|----------|--------------------|------------|----------|-------------------|----------|----------|----------|----------|-------------------|----------|----------|----------|----------|----------|----------|----------|----------|----------|----------|----------|------------|------------|----------|--------------------|-------------------|-------------------|-----|----|----------|------|-------------------|
| 40 | 40 | 40 | 40 | 40 | 40 | 40 | 40 | 40 | 40 | 40 | 40 | 40 | 40 | 40 | 40 | 40 | 40 | 40 | 40 | 40 | 40 | 40 | 40 | 40 | 40 | 40 | 40 | 40 | 40 | 40 | 40 | 40 | 40 | 40 | 40 |
| 0 | 0 | | 0 | .0 | | .0 | | | | | ٥. | | | | | ٥. | .0 | | | | | | | | | | | | | | | | | | 100.0 |
| 745 | 552 | 220 | 36 | 36 | 28 | 28 | 28 | 22 | 22 | 22 | 22 | 22 | 22 | 18 | 18 | 18 | 18 | 17 | 17 | 17 | 17 | 13 | 13 | 13 | <u>1</u> 3 | <u>1</u> 1 | 片 | 1 | 11 | 10 | 10 | 10 | 9 | φ | φ |
| 19 | 21 | 22 | 24 | 23 | 24 | 23 | 23 | 23 | 23 | 23 | 23 | 23 | 23 | 23 | 23 | 23 | 23 | 23 | 23 | 23 | 23 | 23 | 23 | 23 | 23 | 23 | 23 | 23 | 23 | 23 | 23 | 23 | 23 | 23 | 23 |
| AAW49096 | AAY84883 | AAB94488 | ABU08436 | AAM48652 | ABU08434 | AAM48523 | ABB08740 | AAM48637 | AAM48636 | AAM48635 | AAM48634 | AAM48631 | AAM48630 | AAM48633 | AAM48632 | AAM48629 | AAM48628 | AAM48644 | AAM48643 | AAM48639 | AAM48638 | AAM48645 | AAM48642 | AAM48641 | AAM48640 | AAM48653 | AAM48525 | AAM48506 | ABB77311 | 53 | N | 31 | AAM48533 | 853 | AAM48529 |
| | | | | | | | | | | | | | | | | | | | | | , | | | | | | | | | | | | | | • |
| Human I-kappa-B ki | -каррав | m | Human IkappaB kina | tate | Wild-type human NE | eptide SEQ |) bin | Anti-inflammatory | -inf | ١. | 1 | -in | Anti-inflammatory | i-inf | • | -inf | -inf | -inf | i-inf | inf | inf | inf | i-infl | inflamma | ā | peptide. | a | Human IKKbeta pept | Human NBD peptide | Anti-inflammatory | - (| Œ | i-inf | -inf | Anti-inflammatory |

ALIGNMENTS

| DE | ğ | ဌ | ğ | AC | × | ij | ABB087 | RESU |
|-----------------------------|---|---------------|---|-----------|---|--------------------|---------------|--------|
| IKKbeta NEMO | | 14-JUN-2002 | | ABB08725; | | ABB08725 star | 8725 | SULT 1 |
| IKKbeta NEMO binding domain | | (first entry) | | | | standard; peptide; | | |
| <u> </u> | | _ | | | | - | | |
| peptide | | | | | | 6 AA. | | |
| SEQ | | | | | | | | |
| 벙 | | | | | | | | |
| ID NO | | | | | | | | |

IKKbeat; IKKalpha; NEMO; NEMO binding domain; NBD; NF-kappaB; NF-kB; kinase activation; leukocyte; inflammation; E-selectin; osteoclast; autoimmune disease; transplant rejection; osteoporosis; cancer; Alzheimer's disease; viral; infection; astima; anaphylaxis; psoriasis; rheumatoid arthritis; Crohn's disease; multiple sclerosis; HIV; corticosteroid; immunosuppression; antiinflammatory; immunosuppressive; osteopathic; ostostatic; nootropic; neuroprotective; anti-HIV; human; antiarteriosolerotic; virucide; antiasthmatic; antiallergic; dermatological; antibacterial; antipporiatic; antirheumatic; 02-MAY-2001; 2001WO-US40654. 08-NOV-2001 WO200183547-A2 Homo sapiens. antiarthritic; osteopathic; antiulcer

02-MAY-2000; 2000US-201261P 22-AUG-2000; 2000US-0643260

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RESULT 2
AAM48530
ID AAM4
XX
XX
AAC
AAM4
XX
DT 20-M
DT 20-M
DX
DX
ANTI
XX
ANTI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CC omprises contacting a cell with an anti-inflammatory compound (ABB08725-ABB08742) comprising at least one NEWO binding domain (ABB77313). The compound has acts through selective inhibition of (ABB77313). The compound has acts through selective inhibition of (ABB77313). The compound has acts through selective inhibition of (ABB77313). The compound has acts through selective inhibition of (ABB77313). The compound has acts through selective inhibition of (ABB77313). The compound has acts through selective inhibition of (ABB77313). The compound may also condition results in inhibition of IXKbeta kinase activation and subsequent decreased phosphorylation of IXKbeta kinase activation and condition selection of selection of leukocytes care (Alrectly) by blocking the recruitment of leukocytes care (Alrectly or indirectly) by blocking the recruitment of leukocytes care (Alrectly or indirectly) by blocking the recruitment of leukocytes care (Alrectly or indirectly) is blocking the recruitment of leukocytes care (Alrectly or indirectly) is blocking the recruitment of leukocytes of act (directly or indirectly) by blocking the recruitment of leukocytes care (Alrectly or indirectly) by blocking the recruitment of leukocytes of act (directly or indirectly) by blocking the recruitment of leukocytes care (Alrectly) by blocking the recruitment of leukocytes of antimitiation, where the condition is an inflammatory disorder, an autoimmune disease, where the condition is an inflammatory disorder, an autoimmune disease, the condition is an inflammatory disorder, an arthaution, by down-regulating the care inflammatory disorder is arthauting allergies.

CC treating disease, transplant rejection, osteoporosis, cancer (Alrematicis, Booriasis, psoriasis, psoriasis, psoriasis, psoriasis, psoriasis, psoriasis, recember viral infections of treating anaphylaxis, drug and food sensitivity, contact dermatitis, caused by Epstein-barr, cytomegalovirus or herpes simplex. Other viral infections of including and tor transplants of antiflammatory
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              밁
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local S
Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                мау мл,
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Antiinflammatory; antiasthmatic; cytostatic; antipsoriatic; nootropic; antiheumatic; antiasthritic; osteopathic; antibacterial; virucide; immunosuppressive; dermatological; neuroprotective; antiatherosclerotic; antiallergic; membrane translocation domain; NEMO binding domain; eczema
                                                                                                                                                                            Anti-inflammatory peptide SEQ ID NO 33.
                                                                                                                                                                                                                                                    20-MAR-2002
                                                                                                                                                                                                                                                                                                                                                                                              AAM48530 standard; Peptide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1 LDWSWL 6
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6; Conserv
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Pred. No. 9.3e+05;
; Mismatches 0;
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rheumatoid arthritis; osteoarthritis; inflammatory bowel disease; autoimmune disorder; multiple sclerosis; transplant rejection; osteoporosis; Alzheimer's disease; atherosclerosis; viral infection; ataxia telangiectasia; allergy; anaphylaxis; arthritis.
                                                                                                                                                                                                                                                                                         02-MAY-2000; 2000US-201261P.
22-AUG-2000; 2000US-0643260.
                                                                                                                                                                                                                                                                                                                                                          Synthetic
                                                                                                                                                                                                                                                      мау мл,
                                                                                                                                                                                                                                                                                                               02-MAY-2001; 2001WO-US14346
                                                                                                                                                                                                                                                                                                                                             WO200183554-A2
                                                                                                                                                                                                                                                                                                                                                                                                     cytokine; NFkappaB; IkappaB kinase beta;
                                                                                                                                                                                                                                                                    (PRAE-) PRAECIS PHARM INC (UYYA ) UNIV YALE.
                                                                                                                                                                                                                                                      Ghosh S,
                                                                                                                                                                                                                                                      Findeis MA,
                                                                                                                                                                                                                                                      Phillips K;
                                                                                                                                                                                                                                                                                                                                                                                                IKKbeta; cancer; psorias
flammatory bowel disease;
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Movel antiinflammatory compound comprising membrane translocation domain fused to NEMO binding sequence, useful for blocking nuclear factor kappaB activation, and for treating asthma, lung inflammation, psoriasis

WPI;

Claim 6; Page 61; 88pp; English

The invention relates to an antiinflammatory compound (especially

6-15

cc amino acid residues, fused to a NEMO binding sequence (AAM/8525-AAM/8619). The antiinflammatory compounds have antiasthmatic, cyrostatic, antipsoriatic, antiinflammatory compounds have antiasthmatic, cyrostatic, antipsoriatic, antiinflammatory compounds have antiasthmatic, cyrostatic, antipsoriatic, antiinflammatory compounds have antiasthmatic, corporate activatic, antipsoriatic, antipsoriatic, antiinflammatory compounds have antiasthmatic, compounds act as selective inhibitors of cytokine-mediated NFkappas activation by blocking interaction of Ikappas kinase beta (IKKbeta) at the NEMO binding domain that results in inhibition of IKCbeta kinase compounds are useful for treating inflammatory disorders, e.g. asthma, compounds are useful for treating inflammatory disorders, e.g. asthma, consecuritis; antimmune diseases such as lupus, polymyalgia, scleroderma, concentratis; antiple sclerosis; transplant rejection; osteoporosis; callangiectasia. The compounds are also useful for treating conjunction; and ataxia consecuritis.

Conformation of the sclerosis is transplant rejection; and ataxia consecuritis antiple sclerosis; viral infections; and ataxia consecurities and compounds are also useful for treating anaphylaxis, consecurities. AAM48628-AAM48645), comprising a membrane translocation domain (AAM48620-AMM48627 or AAM48646-AAM48651) which comprises from

Sequence 6 AA;

RESULT 3
AAM48655
ID AAM4
XX
AC AAM4
XX
AC AAM4
XX
XX
XX 밁 S Query Match
Best Local Similarity
Matches 6; Conserv AAM48655; AAM48655 LDWSWL 6 standard; Conservative Peptide; 100.0%; Score 40; Di 100.0%; Pred. No. 9.: tive 0; Mismatches 9.3e+05 띪 23; 0 Length 0 Gaps

0

20-MAR-2002

(first entry)

mutant

peptide

SEQ

IJ ö

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HANNE WANTER STANDER OF THE STANDER 
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                                                                                                                                                                                                                                                                                                                                      amino acid residues, fused to a NEMO binding sequence

(AAM48525-AAM48619). The antiinflammatory compounds have antiasthmatic,

(Cytostatic, antipsoriatic, antiirheumatic, antiarthritic, osteopathic,

(C antibacterial, immunosuppressive, dermatological, neuroprotective,

(C antibacterial, immunosuppressive, dermatological, neuroprotective,

(C compounds act as selective inhibitors of cytokine-mediated NFkappaB

(C compounds act as selective inhibitors of cytokine-mediated NFkappaB

(C compounds act as selective inhibitors of IKappaB kinase beta (IKKbeta) at

(C compounds are useful for treating inflammatory disorders, e.g. asthma,

(C compounds are useful for treating inflammatory disorders, e.g. asthma,

(C consounds are useful for treating inflammatory disorders, e.g. asthma,

(C osteoarthritis, antiammatory bowel disease, sepsis, vasculitis,

(C osteoarthritis, aultiple sclerosis, transplant rejection, osteoporosis,

(C dragor food sensitivity, eczema, dermatitis, sunburn, aging and
                                                                                                                           Matches
                                                                                                                                                       Query Match
Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Antinflammatory, antiasthmatic; cytostatic; antipscriatic; nootropic; antirfeumatic; antiarthritic; osteopathic; antibacterial; virucide; antiarthritic; osteopathic; antibacterial; virucide; immunosuppressive; dermatological; neuroprotective; antiatherosclerotic; antiallergic; membrane translocation domain; NEMO binding domain; eczema; cytokine; NFkappaB; IkappaB kinase beta; IKVbeta; cancer; psoriasis; rheumatoid arthritis; osteoarthritis; inflammatory bowel disease; autolimmune disorder; multiple sclerosis; transplant rejection; osteoporosis; Alzheimer's disease; atherosclerosis; viral infection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The invention relates to an antiinflammatory compound (especially AAM48628-AAM48645), comprising a membrane translocation domain (AAM48620-AAM48627 or AAM48646-AAM48651) which comprises from 6-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 2002-121889/16.
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22-AUG-2000;
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                                                                                                                                                                                                                                                   Sequence
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UNIV YALE.
                                                                                                                        Similarity 6; Conser
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                                                      LDWSWL 6
                                                                                                                                                                                                                                                   6 AA;
                                                                                                                           Conservative
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2000US-0643260.
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Pred. No. 9.3
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RESULT 4 ABU08418

RESULT 5 AAM48534

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LDWSWL 6

Gaps

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Query Match
Best Local S
Matches 6
                                                                                                             NEMO binding domain (NBD) peptides. The NEWO binding domains are found on IkappaB kinase-beta (IKKDeta) and IkappaB kinase-apha (IKKDeta) are useful for modulating nuclear factor-kappaB (NF-kappaB) induction in a cell, where the compounds are capable of blocking the interaction between one or more IKKS such as IKKAlpha or IKKDeta, and NEMO. The antiinflammatory compound further comprises at least one membrane translocation domain. The compounds are useful for treating inflammatory disorders, autoimmune diseases, osteoporosis, cancer, Alzheimer's disease, atherosclerosis, viral infections, Ataxia telangiectasia, and for transplantation detection. The compounds of the invention block NF-kappaB induction by IKK but do not inhibit the basal activity of NF-kappaB. ABU08418-ABU08432 represent human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    IkappaB kinase-beta; IkappaB kinase-alpha; IKKalpha; NF-kappaB; muclear factor-kappaB induction; inflammatory disorder; autoimmune disease; osteoporosis; cancer; Alzheimer's disease; atherosclerosis; viral infection; Ataxia telangiectasia; transplantation detection; immunosuppressive; osteopathic; cytostatic; nootropic; neuroprotective; antiatherosclerotic; virucide; vasotropic; antirheumatic; antiarthritic; mutant; mutein.
                                                                                                                                                                                                                                                                                                                                                                                                   Novel antiinflammatory peptide compounds comprising NEMO binding domain, useful for modulating NF-kappaB induction in a cell and for treating NF-kappaB-mediated inflammation disorders e.g., asthma, psoriasis, vasculitis
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N-PSDB; ABX94269, ABX94270.
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Synthetic.
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                                                                 Sequence
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22-AUG-2000;
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                                                                                                   NBD mutant
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GHOSH
 Similarity 6; Conser
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2000US-0643260
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 Score 40; DB 24;
Pred. No. 9.3e+05;
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                                 Length
     Indels
                                     o,
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20-MAR-2002

(first entry)

peptide

SEQ

ID NO

AAM48534 standard; Peptide; 7

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amino acid residues, fused to a NEMO binding sequence
(AAM48525-AAM48619). The antiinflammatory compounds have antiasthmatic,
cyrostatic, antipsoriatic, antirheumatic, antiarthritic, osteopathic,
antipacterial, immunosuppressive, dermatological, neuroprotective,
compounds act as selective inhibitors of cyrokine-mediated NFkappaB
compounds are blocking interaction of IkappaB kinase beta (IKKbeta) at
the NEMO binding domain that results in inhibition of IKCbeta kinase
compounds are useful for treating inflammatory disorders, e.g. asthma,
compounds are useful for treating inflammatory disorders, e.g. asthma,
compounds are useful for treating inflammatory disorders, e.g. asthma,
compounds are useful for treating inflammatory disorders, e.g. asthma,
compounds are useful for treating inflammatory disorders, e.g. asthma,
compounds are useful for treating inflammatory disorders,
compounds are such as lupus, polymyalgia, scleroderma,
compounds are also useful for treating
compounds are a
  Query Match
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Matches 6; Conser
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            immunosuppressive; dermatological; neuroprotective; antiatherosclerotic; antiallergic; membrane translocation domain; NEMO binding domain; eczema cytokine; NFkappaB; IkappaB kinase beta; IKKbeta; cancer; psoriasis; rheumatoid arthritis; osteoarthritis; inflammatory bowel disease; autoimmune disorder; multiple sclerosis; transplant rejection; osteoporosis; Alzheimer's disease; atherosclerosis; viral infection; ataxia telangiectasia; allergy; anaphylaxis; arthritis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Novel antiinflammatory compound comprising membrane translocation domain fused to NEMO binding sequence, useful for blocking nuclear factor kappaB activation, and for treating asthma, lung inflammation,
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22-AUG-2000; 2000US-0643260.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAM48628-AAM48645), comprising a membrane translocation domain (AAM48620-AAM48627 or AAM48646-AAM48651) which comprises from 6-15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim 6; Page 61; 88pp; English
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                                                                                                           Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           invention relates to an antiinflammatory compound (especially
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                                 Score 40; DB 23;
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(UYYA )
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22-AUG-2000;
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2000US-0643260
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Phillips K
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Novel antiinflammatory compound comprising membrane translocation domain fused to NEWO binding sequence, useful for blocking nuclear factor kappaB activation, and for treating asthma, lung inflammation,

Claim 6; Page 61; , dd88 English.

The invention relates to an antiinflammatory compound (especially CA AAM48628-AAM48645), comprising a membrane translocation domain CC (AAM48620-AAM48627 or AAM48621) which comprises from 6-15 cm and acid residues, fused to a NEMO binding sequence composition acid residues, fused to a NEMO binding sequence composition, antispeoriatic, antisheumatic, antiarthritic, osteopathic, corposition, immunosuppressive, dermatological, neuroprotective, compounds act as selective inhibitors of cytokine-mediated NFKappaB compounds are useful for treating inflammatory disorders, e.g. asthma, compounds are useful for treating inflammatory disorders, e.g. asthma, compounds are useful for treating inflammatory disorders, e.g. asthma, compounds are useful for treating inflammatory disorders, e.g. asthma, control inflammation or cancer, psoriasis, rheumatoid arthritis, autoimmune diseases such as lupus, polymyalgia, scleroderma, compounds are such as such as supus, polymyalgia, scleroderma, compounds are also useful for treating of treating of treating ananhylaxis. pro-inflammatory responses such as allergies, drug or food sensitivity, eczema, dermatitis,

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RESULT 7
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ID AAM4
XX AAM4
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XX 
The invention relates to an antiinflammatory compound (especially. CC AAM48628-AAM48627) comprising a membrane translocation domain CC (AAM48620-AMM48627 or AAM48646-AAM48651) which comprises from 6-15 cm and residues, fused to a NEMO binding sequence (AAM48625-AAM48619). The antiinflammatory compounds have antiasthmatic, crycostatic, antipsoriatic, antiinflammatory compounds have antiasthmatic, crycostatic, antiatherosclerotic, virucide and antiallergic activity. The compounds act as selective inhibitors of cytokine-mediated NPkappaB cryotation by blocking interaction of IkappaB kinase beta (IKKbeta) at the NEMO binding domain that results in inhibition of IKkbeta) at compounds are useful for treating inflammatory disorders, e.g. asthma, compounds are useful for treating inflammatory disorders, e.g. asthma, compounds are useful for treating inflammatory disorders, e.g. asthma, compounds are useful for treating inflammatory disorders, e.g. asthma, conflammatory disorders, e.g. asthma, conflammatory inflammatory disorders, e.g. asthma, conflammatory disorders, e.g. asthma, conflammatory inflammatory disorders, e.g. asthma, conflammatory disorders, e.g. a
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              02-MAY-2001; 2001WO-US14346
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim 6; Page 61; 88pp; English
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Pred. No. 9.3e+05;
Nismatches 0;
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CC I-kappa-B kinase (AAW96158), deletion mutants of it retaining CC I-kappa-B kinase activity and I-kappa-B polypeptides (comprising a CC six residue domain of I-kappa-B containing one of Ser32 and Ser36, CC and a candidate agent) can be used to screen for agents that CC modulate the interaction of an IKK polypeptide to a binding target. CC The modulation of the kinase activity of IKK-alpha forms a method CC for modulating signal transduction involving I-kappa-B in a cell. CC for IKK-alpha polypeptides are useful for generating oligonucleotide CC for modulating polypeptides are useful for generating oligonucleotide CC for modulating nucleic acids. The nucleic acids are useful as CC for alpha-encoding nucleic acids. The nucleic acids are useful as CC for modulating probes and primers. Their diagnostic applications CC include IKK-alpha alleles in clinical and laboratory samples. CC meaning the probes for identifying wild-type and CC mitant IKK-alpha alleles in clinical and laboratory samples. CC concentration/availability of active IKK-alpha.

CC catalytically inactive IKK-alpha mutants suppress NF-kappa-B cativation induced by tissue necrosis factor (TNF), interleukin-1

Disclosure; Page -; 32pp; English.

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence
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01-JUL-1997;
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97US-0887115.
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N.B. The present sequence is not given in the present specification but is derived from the sequence given in AAW96157 as specified.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           autoimmune disorder; multiple sclerosis; transplant rejection; osteoporosis; Alzheimer's disease; atherosclerosis; viral infection; ataxia telangiectasia; allergy; anaphylaxis; arthritis.
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                                                                                                                                                                                                                                                                                                                                                         Novel antiinflammatory compound comprising membrane translocation domain fused to NEWO binding sequence, useful for blocking nuclear factor kappaB activation, and for treating asthma, lung inflammation,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     02-MAY-2000;
22-AUG-2000;
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The invention relates to an antiinflammatory compound (especially AAM48628-AAM48645), comprising a membrane translocation domain (AAM48620-AAM48627 or AAM48646-AAM48651) which comprises from 6-15 amino acid residues, fused to a NEMO binding sequence (AAM48625-AAM48619). The antiinflammatory compounds have antiasthmatic, cytostatic, antipsoriatic, antirheumatic, antiarthritic, osteopathic, antibacterial, immunosuppressive, dermatological, neuroprotective,
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                                                                                                                                                                                                  relates to an antiinflammatory compound (especially
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Pred. No. 9.3e+05;
Mismatches 0;
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domain fused to NEMO factor kappaB activat psoriasis

The invention Claim 6; Page

relates to

an antiinflammatory

compound

(especially

61;

WPI; 2002-121889/16.

Novel antiinflammatory compound comprising membrane translocation domain fused to NEMO binding sequence, useful for blocking nuclear factor kappaB activation, and for treating asthma, lung inflammation

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ARMSULT 10
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22-AUG-2000;
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factor kappaB activation, and psoriasis -

for treating asthma,

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22-AUG-2000; 2000US-0643260
                                                                                                                                                                                                                                                                                                                  Synthetic
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Novel antiinflammatory compound comprising membrane translocation domain fused to NEMO binding sequence, useful for blocking nuclear
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Antibacterial, immunosuppressive, dermatological, neuroprotective, cantibacterial, immunosuppressive, dermatological, neuroprotective, mostropic, antiatherosclerotic, virucide and antiallergic activity. The compounds act as selective inhibitors of cytokine-mediated NFkpppaB compounds act as selective inhibitors of IKappaB kinase beta (IKKbeta) at the NEWO binding domain that results in inhibition of IKCbeta kinase activation by blocking increased phosphorylation of IKappaB. The compounds are useful for treating inflammatory disorders, e.g. asthma, lung inflammation or cancer, paoriasis, rheumatoid arthritis, osteoarthritis, inflammatory bowel disease, segsis, vasculitis, costeoarthritis, inflammatory bowel disease, segsis, vasculitis, bursitis, autoimmune diseases such as lupus, polymyalgia, scleroderma, granulomatosis, multiple sclerosis; transplant rejection; osteoporosis; granulomatosis, and atherosclerosis; viral infections; and araxia telangiactasia. The compounds are also useful for treating transplanticy responses such as allergies, urbicaria, anaphylaxis, are food sensitity eczem dermatitis, sunburn, ading and
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAM48628-AAM48645), comprising a membrane translocation domain (AAM48620-AVM48627 or AAM48646-AAM48651) which comprises from
                                                                                  drug or food sensitivity, eczema, dermatitis, sunburn, aging
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밁 Ş Query Match Best Local Matches w 1 LDWSWL 6 6; Conserv LDWSWL 8 100.0%; ilarity 100.0%; Conservative 0 0 Score 40; DB Pred. No. 9.3 0; Mismatches 9.3e+05; hes 0; DB 23; Length 9; Indels ٥, Gaps

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Sequence

9 AA;

Anti-inflammatory peptide SEQ ID NO 36. AAM48533 standard; Peptide; 20-MAR-2002 AAM48533; (first entry) 9

immunosuppressive; dermatological; neuroprotective; antiatherosclerotic; antiallergic; membrane translocation domain; NEMO binding domain; eczema; cytokine; NFkappaB; IkappaB kinase beta; IKCbeta; cancer; psoriasis; rheumatoid arthritis; osteoarthritis; inflammatory bowel disease; autoimmune disorder; multiple sclerosis; transplant rejection; osteoporosis; Alzheimer's disease; atherosclerosis; viral infection; Antiinflammatory; antiasthmatic; cytostatic; antipsoriatic; nootropic; antirheumatic; antiarthritic; osteopathic; antibacterial; virucide; 02-MAY-2000; 22-AUG-2000; 02-MAY-2001; 2001WO-US14346 WO200183554-A2 Synthetic ataxia telangiectasia; 08-NOV-2001. PRAECIS PHARM INCUNIV YALE. 2000US-201261P 2000US-0643260

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                                                                                                                                                                                                                                                                                                                                   IKKbeat; IKKalpha, NEMO; NEMO binding domain; NBD; NF-kB; kinase activation; leukocyte; inflammation; E-selectin; osteoclast; autoimmune disease; transplant rejection; osteoporosis; cancer; Alzheimer's disease; viral; infection; asthma; anaphylaxis; psoriasis; rheumatoid arthritis; Crohn's disease; multiple sclerosis; HIV; corticosteroid; immunosuppression; antiinflammatory; immunosuppressive; osteopathic; cytostatic; nootropic; neuroprotective; anti-HIV; human; antiarteriosclerotic; virucide; antiasthmatic; antiallergic; dermatological; antibacterial; antipsoriatic; antirheumatic; antirheumatic; antiarthritic; osteopathic; antiulcer.
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Pred. No. 9.3e+05;
0; Mismatches 0;
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RESULT 14
AAM48528
ID AAM48
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AC AAM48
XX
AC AAM48
XX
AC AAM48

AAM48528 standard;

Peptide; 10

A

20-MAR-2002 AAM48528;

(first entry

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Query Match Best Local Similarity

Sequence

10 AA;

Matches

<u>ه</u>

Conservative

100.0%; Score 40; DB 100.0%; Pred. No. 3.9; tive 0; Mismatches

23; 0;

Length 10; Indels

0

Gaps

LDWSWL 6

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comprises contacting a cell with an anti-inflammatory compound
(ABB08725-ABB08742) comprising at least one NEMO binding domain
(C (ABB77313). The compound has acts through selective inhibition of
(C (YAB577313). The compound has acts through selective inhibition of
(C (YAB577313). The compound has acts through selective inhibition of
(C (YAB577313). The compound has acts through selective inhibition of
(C (YAB577313). The compound has acts through selective inhibition of
(C (YAB577313). The compound has acts through selective inhibition of
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(C (YAB577313). The compound inhibition of IKAD6ta kinase activation of
(C (YAB577313). The compound inflammation, by down-regulating
(C (YAB577313). The inflammatory discorder of leukocytes
(C (YAB577313). The inflammatory discorder is asthma, allergies,
(C (YAB577313). The inflammatory discorder may also be dermatitis, inflammatory
(C (YAB577313). The inflammatory discorder may also be dermatitis,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Modulating NF-kappaB induction in a cell, useful for treating e.g. inflammatory disorders, osteoporosis and cancer, comprises contacting cell with an anti-inflammatory compound comprising at least one NEMO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     02-MAY-2000; 2000US-201261P.
22-AUG-2000; 2000US-0643260.
                                                    anti-inflammatory agents. The present sequence is that of the NEW binding domain of IKKObeta. Note: The present sequence is not given in the specification but encoded by the polynucleotide given at GenBank Accession No. AR06 nucleotides 2203-2235.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Example 4; Page -; 82pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 2002-179350/23
                                                                                                                                                                             any application in which controlled and cancer therapy. Also for id immunosuppression in transplants and cancer therapy. Also for id antiinflammatory compounds and for diagnosis of an inflammatory antiinflammatory compounds and for diagnosis of an inflammatory anti-normal compound may be administered alone or in combination with ot the compound may be administered alone or in combination with ot the NE.
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iflammatory disorder.
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a cell

Anti-inflammatory

peptide

SEQ

IJ

ö 31

Antiinflammatory; antiasthmatic; cytostatic; antipsoriatic; nootropic; antiinflammatory; antiasthmatic; cytostatic; antibacterial; virucide; immunosuppressive; dermatological; neuroprotective; antiatherosclerotic; antiallergic; membrane translocation domain; NEMO binding domain; eczema; cytokine; NFKappaB; IkappaB kinase beta; IKKbeta; cancer; psoriasis; rheumatoid arthritis; osteoarthritis; inflammatory bowel disease; autoimmune disorder; multiple sclerosis; transplant rejection; osteoporosis; Alzheimer's disease; atherosclerosis; viral infection; autoimmune disorder; multiple sclerosis; transplant rejection; osteoporosis; Alzheimer's disease; atherosclerosis; viral infection;

telangiectasia; allergy; anaphylaxis; arthritis.

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CC AAM/48628-AAM/48645), comprising a membrane translocation domain CC (AAM/48620-AAM/48627 or AAM/48621) which comprises from 6-15 camino acid residues, fused to a NEWO binding sequence amino acid residues, fused to a NEWO binding sequence CC (AAM/48525-AAM/48619). The antiinflammatory compounds have antiasthmatic, cytostatic, antipsoriatic, antiinflammatory compounds have antiasthmatic, cytostatic, antiatheroscierotic, virucide and antiallergic activity. The compounds act as selective inhibitors of cytokine-mediated NFKappaB compounds act as selective inhibitors of cytokine-mediated NFKappaB compounds act as selective inhibitors of stappaB kinase beta (IKKbeta) at the NEMO binding domain that results in inhibition of IKKbeta kinase compounds are useful for treating inflammatory disorders, e.g. asthma, cc lung inflammation or cancer, psoriasis, rheumatoid arthritis, of the compounds are useful for treating inflammatory disorders, e.g. asthma, cc lung inflammatory bowel disease, septys, vasculitis, correctivitis, autoimmune diseases such as lupus, polymyslia, scleroderma, cc burshitis; autoimmune diseases such as lupus, polymyslia, scleroderma, cc langiectasia. The compounds are also useful for treating creating pro-inflammatory responses such as allergies, urticaria, anaphylaxis, corrections of food sensitivity, eczema, dermatitis, sumburn, aging and
문
                        S
                                                         Matches
                                                                    Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                        The invention relates to an antiinflammatory compound (especially
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Claim 6; Page 61; 88pp; English
                                                                                                                Sequence
N
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                                                                      Similarity
                            LDWSWL 6
                                                                                                                  10
                                                        Conservative
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                                                                       100.
                                                                    .0%;
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                                                        Score 40; DB
Pred. No. 3.9
); Mismatches
                                                                       3.9;
                                                                                    23;
                                                         0
                                                                                    Length 10;
                                                           0
                                                         Gaps
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immunosuppressive; dermatological; neuroprotective; antiatherosclerotic; antiallergic; membrane translocation domain; NEMO binding domain; eczema; cytokine; NrkappaB; IkappaB kinase beta; IKKbeta; cancer; psoriasis; rheumatoid arthritis; osteoarthritis; inflammatory bowel disease; autoimmune disorder; multiple sclerosis; transplant rejection; osteoporosis; Alzheimer's disease; atherosclerosis; viral infection; ataxia telangiectasia; allergy; anaphylaxis; arthritis.
                                                                                                                                                                                                                                                                                                                                                   02-MAY-2000;
22-AUG-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAM48531;
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                                                                                                                                                                                                                                                                                                                          (PRAE-)
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                                                                                                                                                                                                                                                                                                                                                                                                                                    Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Antiinflammatory; antiasthmatic; cytostatic; antipsoriatic; nootropic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Anti-inflammatory peptide SEQ ID
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                                                                                                                                                                                                                                                                                        WPI; 2002-121889/16.
                                                                                                                                                                                                                                                                                                         мау мл,
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                                                                                                                                                                                                                                                                                                                                                                                                 08-NOV-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   antirheumatic;
                                                                                                                                                                                                                                                                                                                           PRAECIS PHARM INC. UNIV YALE.
                                                                                                                                                                                                                                                                                                         Ghosh S,
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2000US-0643260.
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Novel antiinflammatory compound comprising membrane translocation domain fused to NEMO binding sequence, useful for blocking nuclear factor kappaB activation, and for treating asthma, lung inflammation,

Novel antiinflammatory compound comprising membrane translocation domain fused to NEMO binding sequence, useful for blocking nuclear factor kappaB activation, and for treating asthma, lung inflammation,

psoriasis

мау мл,

Ghosh S,

Findeis MA,

Phillips K;

2002-121889/16

(UYYA)

PRAECIS PHARM INC. UNIV YALE.

02-MAY-2000; 2000US-201261P. 22-AUG-2000; 2000US-0643260. 02-MAY-2001; 2001WO-US14346 08-NOV-2001

WO200183554-A2

Claim 6; Page 61; 88pp; English

cc (AAM48525-AAM48619). The antinflammatory compounds have antiasthmatic, corycostatic, antipsoriatic, antirheumatic, antiarthritic, osteopathic, antiartheumatic, antiarthritic, osteopathic, cantibacterial, inmunosuppressive, dermatological, neuroprotective, compounds act as selective inhibitors of cytokine-mediated NFkappaB compounds act as selective inhibitors of cytokine-mediated NFkappaB compounds act as selective inhibitors of cytokine-mediated NFkappaB compounds are useful for treating inflammatory thin inhibition of IKappaB compounds are useful for treating inflammatory disorders, e.g. asthma, compounds are useful for treating inflammatory disorders, e.g. asthma, compounds are useful for treating inflammatory disorders, e.g. asthma, contraitis, autoimmune diseases such as lupus, polymyalgia, scleroderma, compounds are also useful for treating compounds are also useful for treating osteoporosis; calheimar's disease, atherosclerosis; viral infections; and ataxia contraitis and ataxia and contraitis and ataxia and contraitis and ataxia and contraitis and ataxia and contraitis and ataxia, are also useful for treating anaphylaxis, contraitis and sensitivity, eczema, dermatitis, sunburn, aging and contraitivity, eczema, dermatitis, sunburn, aging and The invention relates to an antiinflammatory compound (especially AAM48628-AAM48645), comprising a membrane translocation domain (AAM48620-AAM48627 or AAM48646-AAM48651) which comprises from 6-1 6-15

Sequence 10

Query Match 100.0%; 9
Best Local Similarity 100.0%; 1
Matches 6; Conservative 0; Score 40; DB Pred. No. 3.9 0; Mismatches 3.9; 23; 0 Length 10; Indels 0, Gaps

§ 8 Search completed: February 18, 2004, 14:26:17 Job time : 22.7763 secs

Page 10

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OM protein - protein search, using sw
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Listing first 45 summaries
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Maximum DB
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Perfect score:
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seq length: 2000000000
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Match
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87.531 Million cell updates/sec
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Copyright (c) 1993 - 2004 Compugen
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   probable membrane
                                                                                                                  leucyl-tRNA synthe
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             hypothetical protein Rv2975c - Mycobacterium tuberculosis (strain H37RV)

C;Species: Mycobacterium tuberculosis

C;Abte: 17-Utl-1998 #sequence_revision 17-Jul-1998 #text_change 22-Oct-1999

C;Accession: D70672

R;Cole, S.T.; Brosch, R.; Parkhill, J.; Garnier, T.; Churcher, C.; Harris, D.; Garcier, C.; Harris, D.; Garcier, R.; Davles, R.; Devlin, K.; Feltwell, T.; Gentles, S.; Hamlin, N.; Hol:

Rajandream, M.A.; Rogers, J.; Rutter, S.; Seeger, K.; Skelton, S.; Squares, S.

Nature 393, 537-544, 1998

A;Anthors: Sqares, R.; Sulston, J.E.; Taylor, K.; Whitehead, S.; Barrell, B.G.

A;Anthors: Sqares, R.; Sulston, J.E.; Taylor, K.; Whitehead, S.; Barrell, B.G.

A;Anthors: D0ciphering the biology of Mycobacterium tuberculosis from the complete A;Reference number: A70500; MUID:98295987; PMID:9634230

A;Recession: D70672

A;Status; preliminary; nucleic acid sequence not shown; translation not shown

A;Residues: 1-84 <COL.

A;Residues: 1-84 <COL.
                                                                                                                                                                                                                                                                                                                   RESULT 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   A;Cross-references: GB:Z83018; GB:AL123456; NID:g3261671; PIDN:CAB05437.1; PID:e A;Experimental source: strain H37Rv C;Genetics:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     D70672
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Best Local S
Matches 6
                                                  ;Genome: plasmid pNL1
;Note: orf633
                                                                                    ;Genetics:
   Query Match
Best Local Similarity
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ALIGNMENTS

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R.Romine, M.F.; Stillwell, L.C.; Wong, K.K.; Thurston, S.J.; Sisk, E.C.; Sensen, submitted to the EMBL Data Library, July 1998
A,Description: Complete sequence of a 184 kb catabolic plasmid from Sphingomonas A,Reference number: 220992
A;Accession: T31200
A,Accession: T31200
A,Status: preliminary; translated from GB/EMBL/DDBJ
A,Molecule type: DNA
A,Residues: 1-129 <ROM>
A,Gross-references: EMBL:AF079317; NID:g3378261; PID:g3378341; PIDN:AAD03924.1
                                                                                                                                                                                                                                                                                                                                                           hypothetical protein 633 - Sphingomonas aromaticivorans plasmid pNL1 C;Species: Sphingomonas aromaticivorans C;Date: 11-Jan-2000 #sequence_revision 11-Jan-2000 #text_change 11-Jan-2000 C;Accession: T31200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          100.0%; Score 26; DB ilarity 100.0%; Pred. No. 12; Conservative 0; Mismatches
100.0%;
Score
Pred.
  No.
     19;
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                       Length 129;
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RESULT
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8;Description: Sulfolobus solfataricus complete genome.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 R;Nierman, W.C.; Feldblyum, T.V.; Paulsen, I.T.; Nelson, K.B.; Eisen, J.; Heidelberg, J. B.; Laub, M.T.; DeBoy, R.T.; Dodson, R.J.; Durkin, A.S.; Gwinn, M.L.; Haft, D.H.; Kolon, J.; Ermolaeva, M.; White, O.; Salzberg, S.L.; Shapiro, L.; Venter, J.C.; Fraser, C.M. Proc. Natl. Acad. Sci. U.S.A. 98, 4136-4141, 2001
A;Title: Complete Genome Sequence of Caulobacter crescentus.
A;Reference number: A87249; MUID:21173698; PMID:11259647
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 conserved hypothetical protein [imported] - Sulfolobus solfataricus
C;Species: Sulfolobus solfataricus
C;Date: 24-May-2001 #sequence_revision 24-May-2001 #text_change 24-May-2001
                   AtrA transcription regulator [imported] - Sinorhizobium meliloti (strain 1021) C;Species: Sinorhizobium meliloti C;Species: Sinorhizobium meliloti C;Date: 24-Aug-2001 #sequence_revision 24-Aug-2001 #text_change 30-Sep-2001 C;Accession: 595326 C;Accession: 595326 R;Barnett, M.J.; Fisher, R.F.; Jones, T.; Komp, C.; Abola, A.P.; Barloy-Hubler.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              hypothetical protein CC3064 [imported] - Caulobacter crescentus
C;Species: Caulobacter crescentus
C;Date: 20-Apr-2001 #sequence_revision 20-Apr-2001 #text_change 20-Apr-2001
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A; Residues: 1-130 < KUR>
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A; Accession: F90278
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A;Cross-references: GB:
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A; Residues: 1-171 <S'
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Barnett, M.J.; Fisher, R.F.; Jones, T.; Komp, C.; Kalman, S.; Keating, D.H.; Palm, C.; Peck, M.C.;
                                                                                                                                                                                                                                                                                                             Local
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6; Conserv
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Surzycki, R.; Wells, D.H.;
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A;Cross-references: GB:AE006469; PIDN:AAK65175.1; PID:g14523620; GSPDB:GN00165 A;Experimental source: strain 1021, megaplasmid pSymA A;Experimental source: strain 1021, megaplasmid pSymA R;Galibert, F.; Finan, T.M.; Long, S.R.; Puhler, A.; Abola, P.; Ampe, F.; Barlc pela, D.; Chain, P.; Cowie, A.; Davis, R.W.; Dreano, S.; Federspiel, N.A.; Fish L.; Hyman, R.W.; Jones, T. Science 293, 668-672, 2001
A;Authors: Kahn, D.; Kahn, M.L.; Kalman, S.; Keating, D.H.; Kiss, E.; Komp, C.; hebault, P.; Vandenbol, M.; Vorholter, F.J.; Weidner, S.; Wells, D.H.; Wong, K. A;Title: The composite genome of the legume symbiont Sinorhizobium meliloti. A;Reference number: A96039; MUID:21368234; PMID:11474104
                                                                                                                                                                                                                                                                                                       R;Klenk, H.P.; Clayton, R.A.; Tomb, J.F.; White, O.; Nelson, K.B.; Ketchum, K.A.; Fleischmann, R.D.; Quackenbush, J.; Lee, N.H.; Sutton, G.G.; Gill, S.; Kirknes Glodek, A.; Zhou, L.; Overbeek, R.; Gocayne, J.D.; Weidman, J.F.; McDonald, L. Nature 390; 364-370, 1997

A,Authors: Utterback, T.; Cotton, M.D.; Spriggs, T.; Artiach, P.; Kaine, B.P.; S)
Smith, H.O.; Woese, C.R.; Venter, J.C.
A;Title: The complete genome sequence of the hyperthermophilic, sulfate-reducing A,Reference number: A69250; MUID:98049343; PMID:9389475

A,Accession: F69311
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A;Title: Nucleotide sequence and predicted functions of the entire Sinorhizobium
A;Reference number: A95262; MUID:21396509; PMID:11481432
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 conserved hypothetical protein AF0494 - Archaeoglobus fulgidus
C;Species: Archaeoglobus fulgidus
C;Date: 05-Dec-1997 #sequence_revision 05-Dec-1997 #text_change 21-Jul-2000
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149 LDASAL 154
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S.; Federspiel, N.A.; Fisher
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#text_change 31-Dec-2000

(strain PAO1)

; Warrener, K.R.; Kas,

Hickey, Larbig,

PA01,

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R;Goodner, B.; Hinkle, G.; Gat
A; Liu, F.; Wollam, C.; Alli
Science 294, 233-2328, 2001
A;Title: Genome Sequence of th
A;Reference number: A97359; MU
A;Accession: H98287
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A;Status: preliminary
A;Molecule type: DNA
A;Residuse: 1-281 <STO>
A;Cross-references: GB:AE004447; GB:AE004091; NID:g9945902; PIDN:AAG03476.1; GSPDB:GN001
A;Experimental source: strain PAO1
C;Genetics:
C;Genetics:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               probable DNA-binding regulator - Streptomyces coelicolor
C;Species: Streptomyces coelicolor
C;Date: 03-Dec-1999 #sequence_revision 03-Dec-1999 #text_change 03-Dec-1999
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                                                                                                                                                                                                                                                                                                                                      hypothetical protein AGR_L_2514 [imported] - Agrobacterium tumefaciens (strain C;Species: Agrobacterium tumefaciens C;Date: 22-Oct-2001 #sequence_revision 22-Oct-2001 #text_change 18-Nov-2002 C;Accession: H98287
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A; Accession: T37024
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A;Experimental source: strain A3(2)
                                                                                                                                                                           A;Cross-references: GB:AE007870; PIDN:AAK89826.1; PID:g15159760;
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A; Residues: 1-383 < KUR>
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RESULT 10
H81807
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hypothetical protein Rv2370c - Mycobacterium tuberculosis (strain H37RV)
C;Species: Mycobacterium tuberculosis
C;Species: Nycobacterium tuberculosis
C;Date: 17-Uul-1998 #sequence_revision 17-Jul-1998 #text_change 22-Oct-1999
C;Accession: A70587
R;Cole, S.T; Brosch, R.; Parkhill, J.; Garnier, T.; Churcher, C.; Harris, D.
R;Connor, R.; Davies, R.; Devlin, K.; Feltwell, T.; Gentles, S.; Hamlin, N.;
Rajandream, M.A.; Rogers, J.; Rutter, S.; Seeger, K.; Skelton, S.; Squares, Nature 393, 537-544, 1998
Nature 393, 537-544, 1998
A;Authors: Sqares, R.; Sulston, J.E.; Taylor, K.; Whitehead, S.; Barrell, B.9
A;Title: Deciphering the biology of Mycobacterium tuberculosis from the comp
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A,Reference number: A81775; MUID:20222556; PMID:10761919
A,Accession: H81807
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A70587
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         C;Species: Neisseria meningitidis
C;Date: 31-Mar-2000 #sequence_revision 31-Mar-2000 #text_change 19-Jan-2001
C;Accession: B81062
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A;Residues: 1-394 <PAR>
A;Cross-references: GB:AL162757; GB:AL157959; NID:g7380371;
A;Experimental source: serogroup A, strain Z2491
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          A;Authors: Grandi, G.; Sun, L.; Smith, H.O.; Fraser, C.M.; Moxon, E.R.; A;Title: Complete genome sequence of Neisseria meningitidis serogroup B A;Reference number: A81000; MUID:20175755; PMID:10710307
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A; Residues: 1-394 <TET>
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C;Species: Neisseria meningitidis
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Matches 6
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Holroyd, S., Jagels, K.,
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se: serogroup B, strain MC58
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Mungall, K.; Quail,
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Dougherty
Pizza, M.
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A;Status: preliminary; nucleic acid sequence not shown; translation not shown
A;Molecule type: DNA
A;Residues: 1-437 <COL>
A;Cross-references: GB:Z95208; GB:AL123456; NID:g3261747; PIDN:CAB08469.1; PID:e315159;
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H81847
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A;Title: The genome sequence of the facultative intracellular pathogen Brucella melitens A;Reference number: AD3252; PMID:11756688
A;Recession: AH3265
A;Status: preliminary
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C;Superfamily: fumarate hydratase
C;Keywords: ammonia-lyase; carbon-nitrogen lyase
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                                                                                                                                                       A; Molecule type: DNA
A; Residues: 1-512 < PAR>
A; Cross-references: GB:AL162756; GB:AL157959; NID:g7380091; PIDN:CAB84784.1; PID:g7380194; Experimental source: serogroup A, strain Z2491
3; Genetics:
                                                                                                                                                                                                                                                                                                                                                                                       A;Title: Complete DNA sequence of a serogroup A strain of Neisseria menigitidis Z2491.
A;Reference number: A81775; MUID:20222556; PMID:10761919
A;Accession: H81847
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Pred. No.
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Pred. No. 81;
0; Mismatches
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Search completed: February 18, Job time: 8.5921 secs

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R,Goodner, B.; Hinkle, G.; Gattung, S.; Miller, N.; Blanchard, M.; Qurollo, B.; G. A.; Liu, F.; Wollam, C.; Allinger, M.; Doughty, D.; Scott, C.; Lappas, C.; Marke Science 294, 2323-2328, 2001 les and Biotechnology Agent Agrobacter A,Title: Genome Sequence of the Plant Pathogen and Biotechnology Agent Agrobacter A,Reference number: A97359; MUID:21608551; PMID:11743194
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                                                                                                                                      A; Map position: linear c
C; Superfamily: response
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A;Molecule type: DNA
A;Residues: 1-513 <KUR>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   RESULT
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3: sp_funga:*
4: sp_human:*
5: sp_invertebrate:*
6: sp_mammal:*
7: sp_mhc:*
9: sp_organelle:*
9: sp_phage:*
1: sp_vodent:*
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1: sp_virus:*
1: sp_unclassified:*
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Q97ys3 sulfolobus
Q96zni sulfolobus
Q9a3y6 caulobacter
Q8k1x4 pseudomonas
Q9ky23 streptomyce
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Q9ck50 streptomyce
Q9zgg rhizobium m
Q94iz7 oryza sativ
Q9hfu7 pneumocysti
Q9xcy6 vibrio para
Q92756 archaegglob
Q91746 pseudomonas
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SPECIES-O.sativa (japonica cultivar-group); STRAIN=cv. Nipponbare;
Sasaki T., Matsumoto T., Yamamoto K.;
Sasaki T., Matsumoto T., Yamamoto K.;
Chromosome 1, BAC

"Oryva sativa nipponbare(GA3) genomic DNA, chromosome 1, BAC

clone:OJ1276_B06.",
Clone:OJ1276_B06.",
Clone:OJ1276_B06.",
Submitted (FEB-2001) to the EMBL/GenBank/DDBJ databases.

EMBL; APO02912; BAB39922.1;
EMBL; APO02912; BAB39925.13.1;

EMBL; APO03339; BAB92513.1;

EMBL; APO03339; BAB92513.1;

SEQUENCE 92 AA; 9407 MW; 6DE88E32F046CD92 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    01-JUN-2001 (TrEMBLrel. 17, Created)
01-JUN-2001 (TrEMBLrel. 21, Last sequence update)
01-JUN-2001 (TrEMBLrel. 22, Last annotation update)
01-OCT-2002 (TrEMBLrel. 22, Last annotation update)
01-OCT-2002 (TrEMBLrel. 23, Last annotation update)
00028E10.27 protein (OJ1276_B06.22 protein).
01028E10.27 OR OJ1276_B06.22.
010228 sativa (Fide), and
01028 sativa (Fide)
                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE FROM N.A.

SPECIES=C. sativa; STRAIN=cv. Nipponbare;
Sasaki T., Matsumoto T., Yamamoto K.;
Sasaki T., Matsumoto T., Yamamoto K.;
"Oryza sativa nipponbare(GA3) genomic DNA, chromosome 1,
clone:P0028E10.";
Submitted (NOV-2000) to the EMBL/GenBank/DDBJ databases.
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         1 LDASAL 6
                                                 Similarity 6; Conserv
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QBnrb3 corynebacte
QBpB54 vanthomonas
Q9v9c6 drosophila
QBvtf1 bacillus st
Q9v16 bacillus st
Q9v167 bacillus st
Q9v197 arabidopsis
QB197 arabidopsis
QB199 arabidopsis
QB190 cryza sativ
Q9eqm9 rattus norv
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Q8yjh4 brucella me
Q8fyc6 brucella me
Q8fyc6 brucella su
Q9ju05 neisseria m
Q8u994 agrobacteri
Q9jz08 neisseria m
Q9jz08 neisseria m
Q9jz08 neisseria m
Q9jz08 arabidopsis
Q9jly47 arabidopsis
Q9nle7 corynebacte
Q9fy94 zea mays (m
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Q9ri53 streptomyce
Q9vr43 drosophila
Q8u4w2 agrobacteri
Q9jye3 neisseria m
Q9jye3 neisseria m
Q9jf41 neisseria m
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P95120;
01-MAY-1997 (TrEMBLrel. 03, Created)
01-MAR-2002 (TrEMBLrel. 20, Last sec
01-MAR-2003 (TREMBLREL 23, Last and
Hypothetical procein Rv2975c.
Rv2975C OR MT3052.1 OR MTCY349.12.
                                                                            O85909;
O85909;
O1-NOV-1998
O1-NOV-1998
O1-OCT-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                MEDLINE-9825987; PubMed=9634230;
Cole S.T., Brosch R., Parkhill J., Garnier T., Churcher C., Harris Gordon S.V., Eiglmeier K., Gas S., Barry C.E. III, Tekaia F., Gordon S.V., Eiglmeier K., Gas S., Barry C.E. III, Tekaia F., Badcock K., Basham D., Brown D., Chillingworth T., Connor R., Bavies R., Devlin K., Feltwell T., Gentles S., Hamlin N., Holroyd Hornsby T., Jagels K., Krogh A., McLean J., Moule S., Murphy L., Oliver S., Osborne J., Quail M.A., Rajandream M.A., Rogers J., Rutter S., Seeger K., Skelton S., Squares S., Squares R., Sulston J.E., Taylor K., Whitehead S., Barrell B.G.; "Deciphering the biology of Mycobacterium tuberculosis from the complete genome sequence.";
Nature 393:537-544(1998)."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mycobacterium tuberculosis.
Bacteria; Actinobacteria; Actinobacteridae; Actinomycetales;
Corynebacterineae; Mycobacteriaceae; Mycobacterium.
NCBI_TaxID=1773;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE FROM N.A.
STRAIN=CDC 1551 / Oshkosh;
Fleischmann R.D., Alland D., Eisen J.A., Carpenter L., White O.,
Fleischmann R.D., Alland D., Eisen J.A., Carpenter L., White O.,
Peterson J., DeBoy R., Dodson R., Gwinn M., Haft D., Hickey E.,
Kolonay J.F., Nelson W.C., Umayam L.A., Ermolaeva M., Salzberg S.L.,
Kolonay J.F., Nelson W.C., Umayam L.A., Khouri H., Gill J., Mikula A.,
Delcher A., Utterback T., Weidman J., Khouri H., Gill J., Mikula A.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                  laboratory strains.";
Submitted (APR-2001) to the EMBL/GenBank/DDBJ databases.
EMBL; Z83018; CAB05437.1; ALT_INIT.
EMBL; AB007126; AAK47379.1; -.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                STRAIN=H37RV;
MEDLINE=98295987;
                                              Hypothetical ORF633.
                                                                                                                                                                                                                                                                                                                                                    InterPro; IPR001969; Aspprotease site.
PROSTIE; PS00141; ASP PROTEASE; I.
Hypothetical protein; Complete proteome.
SEQUENCE 92 AA; 9850 MW; 50BDIAFCFCDFD253 CRC64;
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TubercuList; Rv2975c; -
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                               Sphingomonas aromaticivorans
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Proteobacteria; Alphaproteobacteria; Sphingomonadales;
                                                              (TrEMBLrel. 08, Created)
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Pred. No.
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RESULT 5
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RESULT 1037183 CONTROL TO THE PROPERTY OF THE 
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InterPro; IPR002715; PIN.
InterPro; IPR006596; PIN.
InterPro; IPR006596; PIN.
Pfam; PF01850; PIN; 1.
SMANT; SM00670; PIN; 1.
SMANT; Complete p:
Hypothetical protein; Complete p:
SEQUENCE 130 AA; 15118 MW; 1.
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01-OCT-2001
01-OCT-2001
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STRAIN-ATCC 35092 / DSM 1617 / P2;

MEDLINE=21332296; PubMed=11427726;

MEDLINE=21332296; PubMed=11427726;

She Q., Singh R.K., Confalonieri F., Zivanovic Y., Allard G.,

She Q., Singh R.K., Confalonieri F., Zivanovic Y., Allard G.,

Awayez M.J., Chan-Weiher C.C.-Y., Clausen I.G., Curtis B.A.,

De Moors A., Erauso G., Fletcher C., Gordon P.M.K.,

Heikamp-de Jong I., Jeffries A.C., Kozara C.J., Medina N., Peng X.,

Thi-Ngoc H.P., Redder P., Schenk M.E., Theriault C., Tolstrup N.,

Charlebois R.L., Doolittle W.F., Dunk, Gaasterland T.,

Garrett R.A., Ragan M.A., Sensen C.W., Van der Oost J.;

"The complete genome of the crenarchaeon Sulfolobus solfatarious P2.";

Proc. Natl. Acad. Sci. U.S.A. 98:7835-7840(2001).
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NCBI_TaxID=48935;
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(TrEMBLrel. 18, Last sequence update)
(TrEMBLrel. 23, Last annotation update)
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Q9A3Y6;
01-JUN-2001
01-JUN-2001
01-MAR-2002
                                                                                                                                                                                                                                                                   Nierman W.C., Feldblyum T.V., Laub M.T., Paulsen I.T., Nelson Elsen J., Heidelberg J.F., Alley M.R.K., Ohta N., Maddock J.R. Potocka I., Nelson W.C., Newton A., Stephens C., Phadke N.D., Deboy R.T., Dodson R.J., Durkin A.S., Gwinn M.L., Haft D.H., Kolonay J.F., Smit J., Craven M.B., Khouri H., Shetty J., Berr Uterback T., Tran K., Wolf A., Vamathevan M., Ermolaeva M., W Salzberg S.L., Venter J.C., Shapiro L., Fraser C.M.; "Complete genome sequence of Caulobacter crescentus."; Proc. Natl. Acad. Sci. U.S.A. 98:4136-4141(2001).
EMBL, AE005969; AAX25026.1; -.
ETIGR; CC3064; -.
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DNA Res. 8:123-140(2001).
EMBL; AP000987, BAB66893.1;
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STRAIN=ATCC 19089 /
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STRAIN=JCM 10545 / 7;
PubMed=11572479;
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                                                                                                                                                                                                                   Hypothetical SEQUENCE 1
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MEDLINE=21173698; PubMed=11259647;
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01-OCT-2000 (TrEMBLrel. 15, Created)
01-OCT-2000 (TrEMBLrel. 15, Last sequence update)
01-MAR-2003 (TrEMBLrel. 23, Last annotation update)
Hypothetical protein SC02367.
SC02367 OR SCC8A.25C.
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   SEQUENCE FROM N.A.
STRAIN=A3(2);
MEDLINE=97000351; PubMed=8843436;
MEDLINE=97000351; PubMed=8843436;
Redenbach M., Kieser H.M., Denapaite D., Eichner A.,
Rinashi H., Hopwood D.A.;
"A set of ordered cosmids and a detailed genetic and
the 8 Mb Streptomyces coelicolor A3(2) chromosome.";
Mol. Microbiol. 21:77-96(1998).
[4]
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Streptomycineae;
NCBI_TaxID=1902;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Streptomyces coelicolor.

Bacteria; Actinobacteria; Actinobacteridae; Actinomycetales; Streptomycineae; Streptomycetaceae; Streptomyces.
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InterPro; IPR001450; 4Fe45 ferredoxin.
Pfam; PF00037; fer4; 1.
Pfam; PF04060; FeS; 1.
PROSITE; PS00198; 4FE45 FERREDOXIN; 2.
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01-MAR-2003
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01-OCT-2002
                                                                                                                                                                                                                                                                                                      SEQUENCE FROM N.A.
STRAIN=A3(2);
Bentley S.D., Parkhill i
Submitted (MAY-2000) to
                                                                                                                                                                                                                                                                                                                                                                                                                                                             STRAIN=A3(2);
Brown S.P., Harris D
Submitted (MAY-2000)
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Bacteria; Proteobacteria; Gammaproteobacteria; Pseudomonadales;
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E 174 AA; 17648 MW;
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genes in Pseudomonas
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EMBL/GenBank/DDBJ databases.
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annotation update)
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Nature 417:141-147(2002).
EMBL; AL939112; CAB92843.1; -.
InterPro; IPR003325; TerD.
Pfam; PF02342; TerD; 1.
Hypothetical protein; Complete pi SEQUENCE 191 AA; 20414 MW; C.
SEQUENCE From M.145;
STRAIN-A3(2) / M.145;
MEDLINE-21996410; PubMed=12000953;
Bentley S.D., Chater K.F., Cerdeno-Tarraga A.-M., Challis G.L.,
Thomson N.R., James K.D., Harris D.E., Quail M.A., Kieser H.,
Thomson N.R., James K.D., Harris D.E., Quail M.A., Kieser H.,
Cronin A., Fraser A., Goble A., Hidalgo J., Hornsby T., Howarth
Cronin A., Fraser A., Goble A., Hidalgo J., Hornsby T., Howarth
Huang C.-H., Kieser T., Larke L., Murphy L., Oliver K., O'Neil S
Rabbinowitsch E., Rajandream M.A., Rutherford K., Rutter S.,
Rabbinowitsch E., Rajandream M.A., Rutherford K., Rutter S.,
Squares R., Squares R., Squares R., Squares S., Taylor
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Q9KY22;
01-OCT-2000
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               01-OCT-2000 (TrEMBLrel. 15, Created)
01-OCT-2000 (TrEMBLrel. 23, Last sequence update)
01-MAR-2003 (TrEMBLrel. 23, Last annotation update)
Hypothetical protein SC02368.
SC02368 OR SC08A.26C.
Streptomyces coelicolor.
Bacteria; Actinobacteria; Actinobacteria; Actinomycetales;
Streptomycineae; Streptomycetaceae; Streptomyces.
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Bentley S.D.,
Submitted (MAY
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STRAIN=A3(2) / M145;
MEDLINE=21996410; Pu
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                                                                                                                                                                                                                                                                                                                                                                                                                                                         MEDLINE=97000351; PubMed=8843436; Redenbach M., Kieser H.M., Denapaite D.,
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BI_TaxID=1902;
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Nature 417:141-147(2002).
EMBL; AL939112; CAB92844.1;
                                                                                                                                                                                                                                                                                                                                                              STRAIN=A3(2) / M145;

MEDLINE=2196410; PubMed=12000953;

Bentley S.D., Chater K.F., Cerdeno-Tarraga A.-M., Challis G.L.,

Bentley S.D., Chater K.D., Harris D.E., Quail M.A., Kieser H.,

Thomson N.R., James K.D., Harris D.E., Quail M.A., Kieser H.,

Thomson N.R., James K.D., Harris D.E., Chen C.W., Collins M.,

Harper D., Bateman A., Brown S., Chandra G., Chen C.W., Collins M.,

Cronin A., Fraser A., Goble A., Hidalgo J., Hornsby T., Howarth S.,

Huang C.-H., Kieser T., Larke L., Murphy L., Oliver K., O'Neil S.,

Huang C.-H., Kieser T., Larke L., Murphy L., Oliver K., O'Neil S.,

Rabbinowitsch E., Rajandream M.A., Rutherford K., Rutter S.,

Seeger K., Saunders D., Sharp S., Squares R., Squares S., Taylor K.,

Marren T., Wietzorrek A., Woodward J., Barrell B.G., Parkhill J.,

Marren T., Wietzorrek A., Woodward J., Barrell B.G., Parkhill J.,

Marren T., Wietzorrek A., Woodward J., Barrell B.G., Parkhill J.,
                                                 092ZG8 PRELIMINARY;
092ZG8; PRELIMINARY;
01-DEC-2001 (TrEMBLrel. 19,
01-DEC-2001 (TrEMBLrel. 19,
01-MAR-2003 (TrEMBLrel. 23,
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Bacteria; Actinobacteridae; Actinomycetales;
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SCO0641 OR SCF56.25 OR SCF91.01
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01-MAR-2003 (TrEMBLrel. 23, Last
01-MAR-2003 (TrEMBLrel. 23, Last
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Pfam; PF02342; TerD; 1.

Hypothetical protein; Complete
SEQUENCE 191 AA; 20387 MW;
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Nature 417:141-147(2002).
EMBL; AL339105; CAD55270.1; -.
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Pred. No. 1.4e+02;
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39E9C1EC8C47AA7E
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Atra transcriptional regul ATRA OR RA0517 OR SMA0955

Rhizobium meliloti Plasmid pSymA (mega

iloti (Sinorhizobium
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RESULT 12
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ID Q941Z
AC 0941Z
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DT 01-DE
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DE OSUNE
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OC SPERM
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RA SASAK
RT Clone
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AB Barnett M.J., Fisher R.F., Jones T., Komp C., Abola A.P.,
AB Barloy-Hubler F., Boweer L., Capela D., Galibert F., Gouzy J.,
AA Gurjal M., Hong A., Huizar L., Hyman R.W., Kahn D., Kahn M.L.,
AX Kalman S., Keating D.H., Palm C., Peck M.C., Surzycki R., Wells D.H.,
AYeh K.-C., Davis R.W., Federspiel N.A., Long S.R.;
AT "Nucleotide sequence and predicted functions of the entire
sinorhizobium mellioti pSymA megaplasmid.";
Proc. Natl. Acad. Sci. U.S.A. 98:9883-9888(2001).
BR EMBL, AE007243; AAK65175.1; -.
BR InterPro; IPR000524; HTH GNTR, 1.
BR InterPro; IPR000524; HTH GNTR, 1.
BR PAGNITE; PS00043; HTH GNTR, 1.
BR PAGNITE; PS00043; HTH GNTR, 1.
BR PROSITE; PS00043; HTH GNTR, 1.
BR PROSITE; PS00043; HTH GNTR, 1.
BR PROSITE; PS00043; AZ 25843 MW; 82DECAC87E91B94E CRC64;
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Matches 6
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Q94IZ7;
Q94IZ7;
01-DEC-2001 (TrEMBLrel. 19, Created)
01-DEC-2001 (TrEMBLrel. 19, Last sequence update)
01-MR-2003 (TrEMBLrel. 23, Last annotation update)
05JNBa0038J17.13 protein.
05JNBa0038J17.13 protein.
0ryza sativa (Rice).
Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; Liliopsida; Poales; Poaceae;
Ehrhartoideae; Oryzeae; Oryza.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         "Oryza sativa nipponbare(GA3) genomic DNA, chromosome 1, clone:OSJNBa0038017.";
Submitted (JAN-2001) to the EMBL/GenBank/DDBJ databases.
-!- SIMILARITY: CONTAINS 1 RING-TYPE ZINC FINGER.

GRAMENE: AP003104; BAB55721.1; -.
Gramene: Q94127; -.
Interpro. Throcom.
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STRAIN=cv. Nipponbare;
Sasaki T., Matsumoto T., Yamamoto
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STRAIN=1021;
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                                                                                                                                                                                                                                                                                                                        InterPro; IPR001841; Znf_ring.
Pf4am; PF00097; zf_C3HC4; 1.
SMART; SM00184; RING; 1.
PROSITE; PSS0089; ZF_RING_2; 1.
Metal-binding; Zinc; Zinc-finger.
SEQUENCE 237 AA; 23871 MW; EA
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Pred. No. 1.8e+02;
0; Mismatches 0;
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Pred. No.
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Circuit.";

Nucleic Acids Res. 28:0-0(2000).

Nucleic Acids Res. 28:0-0(2000).

EMBL; AF291574; AAG16234.1; -.

EMBL; AF291574; AAG16234.1; -.

EMBL; AF291574; AAG16234.1; -.
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Ivanov I.P., Gesteland R.F., Atkins J.F.;

"Antizyme expression: a subversion of trip

remarkably conserved by evolution, is a si
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     01-MAR-2001 (TrEMBLrel.
01-MAR-2001 (TrEMBLrel.
01-MAR-2001 (TrEMBLrel.
                                                                                                                                                                                                                         J. Bacteriol. 181:3594-3598 (1999).
EMBL, AF119047, AAD39909.1, -.
InterPro, IPR003538, TonB.
InterPro, IPR006260; TonB C.
PRINTS; PR01374; TONBPROTEIN.
                                                                                                                                                                                                                                                                                                                                                            MEDLINE=99287846; PubMed=10348876;
O'Malley S.M., Mouton S.L., Cochino D.A., Deanda
Fuson K.L., Rashidi C.E., Mora M.Y., Payne S.M.,
"Comparison of the heme iron utilization systems
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Vibrio parahaemolyticus.
Bacteria; Proteobacteria;
Vibrionaceae; Vibrio.
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01-NOV-1999 (TrEMBLrel.
01-MAR-2003 (TrEMBLrel.
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SEQUENCE 247 AA; 27121 MW; D949
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NCBI_TaxID=670;
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Pred. No. 1.9e+02;
Mismatches 0;
                                                                                       Score 26; DB 2; 1
Pred. No. 1.9e+02;
; Mismatches 0;
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Henderson D.P.;
of pathogenic
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SETAINN'C-16 / DSP MASSE-389475;

XX MEDLINE=98049343; Pubmed=9389475;

XX Klenk H.-P., Clayton R.A., Tomb J.-F., White O., Nelson K.E.,

XX Klenk H.-P., Clayton R.A., Tomb J.-F., White O., Nelson J.D.,

XX Klenk H.-P., Clayton R.A., Tomb J.-F., White O., Nelson J.D.,

XX Klenk H.-P., Clayton R.J., Gwinn M., Hickey E.K., Peterson J. R.

XX Klenk H.-P., Clayton R.J., Gwinn M., Hickey E.K., Peterson J. Reich C.I., McKenney K., Adams M.D., Loftus B.,

XX Kirkness E.F., Dougherty B.A., McKenney K., Adams M.D., Loftus B.,

XX Kirkness E.F., Dougherty B.A., McKenney T.J., Goldek A., Zhou L.,

RA Kirkness E.F., Dougherty B.A., McKenney J.H., Glodek A., Zhou L.,

RA Cverbeek R., Gocayne J.D., Weidman J.F., McDonald L., Utterback T.,

RA Cotton M.D., Spriggs T., Artiach P., Kaine B.P., Sykes S.M.,

XA Gotton M.D., Spriggs T., Artiach P., Kaine B.P., Sykes S.M.,

XA Asson T.M., Olsen G.J., Fraser C.M., Smith H.O., Woese C.R.,

Wenter J.C.,

"The complete genome sequence of the hyperthermophilic, sulphate-

Treducing archaeon Archaeoglobus fulgidus.";

Nature 390:364-370(1997)

Laboratory A. Argonoff 1: -.
                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local Similarity
Matches 6; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Archaeoglobus fulgidus.
Archaea; Euryarchaeota; Archaeoglobi; Archaeoglobales;
Archaeoglobaceae; Archaeoglobus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       EMBL; AE001070; AAB90743:1; -.
TIGR; AF0494; -.
InterPro; IPR001247; 3 EXCRNASE.
Pfam; PF01138; RNASE PH; 1.
Pfam; PF03725; RNASE PH C; 1.
Hypothetical protein; Complete proteome.
SEQUENCE 259 AA; 28646 MW; E8289D46F9DDCCB3 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NCBI_TaxID=2234;
                                                                                                                                                                                                     149 LDASAL 154
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OM protein -
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seq length: 2000000000
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41.814 Million cell updates/sec
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1: /SIDS1/gcgdata/geneseq/geneseqp-emb1/AA1980.DAT:*

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Gapop 10.0 ,
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GenCore version 5.1.6
Copyright (c) 1993 - 2004 Compugen Ltd.
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Pred. No. is the number of results predicted by chance to hav score greater than or equal to the score of the result being and is derived by analysis of the total score distribution. printed,

SUMMARIES

| Mycobacterium spec | AAY04951 | 20 | 102 | 100.0 | 26 | 9 |
|--------------------|----------|------|--------------------------|-------|-------|---------------|
| Mycobacterium spec | AAY04947 | 20 | 92 | 100.0 | 26 | 80 |
| Mycobacterium spec | AAY04950 | 20 | 84 | 100.0 | 26 | 7 |
| Human mutant NEMO | ABU08435 | 24 | 28 | 100.0 | 26 | Φ |
| NBD peptide SEQ ID | AAM48524 | 23 | 28 | 100.0 | 26 | υ |
| Mutated IKKbeta NE | ABB08741 | 23 | 28 | 100.0 | 26 | 4, |
| Human NEMO binding | ABU08419 | 24 | თ | 100.0 | 26 | ω |
| NBD mutant peptice | AAM48508 | 23 | 0 | 100.0 | 26 | N |
| Mutated IKKbeta NE | ABB08726 | 23 | 6 | 100.0 | 26 | ם |
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| r V |) N |) I | S | 20 | 26 | 26 | 26 | 26 | 26 | 26 | .26 | 26 | 26 | 26 | 26 | 26 | 20 | 26 | 26 | 26 | 26 | 26 | 26 | 26 | 26 | 26 | 26 | 20 | 26 | 26 | 80 | 26 | 26 | 26 | 26 | 26 |
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| 1 | 100 | 2 1 | 967 | 941 | 813 | 745 | 745 | 745 | 745 | 745 | 745 | 669 | 659 | 638 | 600 | 600 | 600 | 600 | 577 | 572 | 564 | 528 | 520 | 516 | 516 | 516 | 462 | 393 | 349 | 284 | 271 | 240 | 160 | 143 | 138 | 102 |
| |) L | ں د | 22 | 23 | 22 | 23 | 22 | 22 | 22 | 22 | 22 | 22 | 22 | 24 | 22 | 22 | 22 | 22 | 21 | 20 | 22 | 22 | 24 | 21 | 21 | 21 | 21 | 24 | 22 | 21 | 21 | 22 | 21 | 21 | 22 | 21 |
| , | 20000000 | | | ABB93761 | | AAM51073 | | AAB79990 | AAB79746 | AAG91007 | AAU71914 | AAE10335 | AAG93061 | AA016363 | AAB80012 | AAB79991 | AAB79747 | AAU71915 | AAG41648 | AAY04954 | AAG41649 | AAU41302 | ABP78709 | AAY75183 | AAY75182 | AAY75180 | AAY75181 | ABP77254 | ABB70854 | AAG51024 | AAG51025 | 9058 | 3553 | വ | 0190 | AAG35536 |
| 10 to | ი. ' | . cinnamoner | Propionibacterium | Herbicidally activ | Drosophila melanog | Corynebacterium gl | Corynebacterium gi | Corynebacterium gi | Corynebacterium gl | glutamicum | C. glutamicum meta | uman transporte | C glutamicum proce | uman epithelial | | | | C. glutamicum meta | is thali | cobacterium spe | is thall | terium | N. gonorrhoeae ami | eisseria menir | isseria | isseria | isseria | N. gonorrhoeae ami | Drosophila melanog | is th | s thaii | nm prot | ģ | sis thali | peptid | tha |

ALIGNMENTS

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          IKKbeat; IKKalpha; NEMO; NEMO binding domain; NBD; NF-kappaB; NF-kB; kinase activation; leukocyte; inflammation; B-selectin; osteoclast; autoimmune disease; transplant rejection; osteoporosis; cancer; Alzheimer's disease; viral; infection; asthoma; anabylaxis; psoriasis; rheumatoid arthritis; Crohm's disease; multiple sclerosis; HIV; corticosteroid; immunosuppression; antiinflammatory; immunosuppressive; osteopathic; cytostatic; mootropic; neuroprotective; anti-HIV; human; antiarteriosclerotic; virucide; antiasthmatic; antiallergic; dermatological; antibacterial; antipsoriatic; antirheumatic; antiarthritic; osteopathic; antiulcer; mutant; mutein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                Homo sapiens.
Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mutated IKKbeta NEMO binding domain peptide SEQ ID NO 3.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              14-JUN-2002 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ABB08726;
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                                                                                                                                                                                                            Misc-difference
                                                                                                                                                                                                                                                                                                                     Misc-difference
                                                                                                                                                                                                                                                                                                                           Location/Qualifiers
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/note= "Wildtype Trp substituted by Ala"
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RESULT 2
AAM48508
ID AAM4
XX
AC AAM4
XX
DT 20-N
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CC comprises contracting a cell with an anti-inflammatory compound (CC (ABB08725-ABB08742) comprising at least one NEMO binding domain (CC (ABB77313). The compound has acts through selective inhibition of (CC (Ybokine-mediated NF-KB activation by blocking the interaction of NEMO CC with IKKbeta at the NEMO binding domain. Blockage of IKKbeta-NEMO CC with IKKbeta at the NEMO binding domain. Blockage of IKKbeta-NEMO CC with IKKbeta at the NEMO binding domain. Blockage of IKKbeta-NEMO CC with IKKbeta at the NEMO binding domain. Blockage of IKKbeta-NEMO CC with IKKbeta at the NEMO binding domain. Blockage of IKKbeta-NEMO CC with IKKbeta at the NEMO binding domain. Blockage of IKKbeta-NEMO CC with IKKbeta at the Compound of IkappaB. The compound may also act (directly or indirectly) by blocking the recruitment of leukocytes of the compound may also conditions, where the condition is an inflammatory disorder, an cc autoimmune disease, atherosclerosis, a viral infection or ataxia conditions, where the condition is an inflammatory disorder, an activation arthritis, ostecoarthritis, psoriatic arthritis, inflammatory classification or ataxia converted arthritis, ostecoarthritis, psoriatic arthritis, inflammatory disorder is asthma, allergies, converted arthritis, psoriatic arthritis, inflammatory converted arthritis, inflammatory disorder may also be dermatitis, eczema, compound anticoarthritis, lupus and converted at the compound may also be dermatitis, converted by Epstein-barr, cytomegalovirus or herpes simplex. Other viral converted by Epstein-barr, cytomegalovirus or herpes simplex. Other viral converted dermatitis, converted d
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              02-MAY-2000;
22-AUG-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 2002-179350/23
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                binding domain -
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                                            AAM48508;
                                                                                                                                                                                                                                                                                                                                                                         Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                 binding
                                                                                                                                                                                                                                                                                                                                                                                                                                                      immunosuppression in transplants and cancer therapy. Also for identifying antiinflammatory compounds and for diagnosis of an inflammatory disorder. The compound may be administered alone or in combination with other known
  20-MAR-2002
                                                                                   AAM48508 standard; Peptide;
                                                                                                                                                                                                                                         1 LDASAL 6
                                                                                                                                                                                                                                                                                   Similarity 6; Conser
                                                                                                                                                                                                                                                                                                                                                                                                                   domain of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Ghosh S;
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2000US-0643260
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                                                                                                                                                                                                                                                                                                                                                                         AA;
    (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                   agents. The present sequence is that of a mutated NEMO IKKbeta.
                                                                                                                                                                                                                                                                                     100.0%; Score 26; DB 23;
100.0%; Pred. No. 9.3e+05;
tive 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            modulating NF-kappaB (NF-kB) induction in ell with an anti-inflammatory compound
                                                                                        6 AA
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RESULT

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1 LDASAL 6

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cc amino acid residues, fused to a NEMO binding sequence (AAM48525-AAM48619). The antiinflammatory compounds have antiasthmatic, cytostatic, antiatherostic, antiinflammatory compounds have antiasthmatic, cytostatic, antiatherostic, antiinflammatory compounds have antiasthmatic, conspounds activatic, immunosuppressive, dermacological, neuroprotective, compounds act as selective inhibitors of cytokine-mediated NFkappaB compounds act as selective inhibitors of cytokine-mediated NFkappaB compounds act as selective inhibitors of cytokine-mediated NFkappaB compounds are useful for treating inflammatory dispates. The compounds are useful for treating inflammatory dispates. The compounds are useful for treating inflammatory dispates, cy asthma, consecuritis, antimune diseases such as lupus, polymyalgia, scleroderma, consecuritis, anticiple sclerosis, transplant rejection, osteoporosis, and ctain confidents are also useful for treating pro-inflammatory responses such as allergies, urticaria, anaphylaxis, consecuritis and consecuritis, activitis, activitis, activitis, activitis, eczema, dermatitis, sunburn, aging and consecuritis are compounds are also useful for treating anaphylaxis, consecuritis and consecuritis activity, eczema, dermatitis, sunburn, aging and
    Matches
                  Query Match
Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             BN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Novel antiinflammatory compound comprising membrane translocation domain fused to NEMO binding sequence, useful for blocking nuclear factor kappaB activation, and for treating asthma, lung inflammation,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Мау МЈ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (PRAE-)
(UYYA )
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        02-MAY-2000; 2000US-201261P
22-AUG-2000; 2000US-0643260
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   02-MAY-2001; 2001WO-US14346
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Antiinflammatory; antiasthmatic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAM48628-AAM48645), comprising a membrane translocation domain (AAM48620-AAM48627 or AAM48646-AAM48651) which comprises from 6-15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     The invention relates to an antiinflammatory compound (especially AAM48628-AAM48645), comprising a membrane translocation domain
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Example 6; Page 47; 88pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    psoriasis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 2002-121889/16
                                                                              Sequence
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l Similarity
6; Conserv
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    Conservative
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    Score 26; DB
Pred. No. 9.3
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    ; DB 23;
. 9.3e+05;
ches 0;
                                          Length 6;
        0
        Gaps
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12-JUN-2003 ABU08419;

(first entry)

kinase activation; leukocyte; inflammation; E-selectin; osteoclast; autoimmune disease; transplant rejection; osteoporosis; cancer; Alzheimer's disease; viral; infection; asthma; anaphylaxis; psoriasis; rheumatoid arthritis; Crohn's disease; multiple sclerosis; HIV; corticosteroid; immunosuppression; antiinflammatory; immunosuppressive; osteopathic; cytostatic; nootropic; neuroprotective; anti-HIV; human; antiarteriosclerotic; virucide; antiasthmatic; antiallergic; antiarteriosclerotic; virucide; antiasthmatic; antiallergic;

Mutated IKKbeta NEMO binding domain peptide SEQ ID NO

NEMO; NEMO binding domain; NBD;

NF-kappaB; NF-kB;

19

IKKbeat; IKKalpha;

14-JUN-2002

(first entry)

ABB08741 standard; peptide; 28

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ABU08419 standard;

peptide;

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ABU08419

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                                                                                                                   Query Match
Best Local Similarity
Matches 6; Conser
                                                                                                                                                                                                                                                                                                                                                                     The present invention relates to antiinflammatory compounds comprising MEMO binding domain (NBD) peptides. The NEMO binding domains are found on IkappaB kinase-beta (IKKbeta) and IkappaB kinase-alpha (IKKalpha) proteins. The antiinflammatory compounds of the invention are useful for modulating nuclear factor-kappaB (NF-kappaB) induction in a cell, where the compounds are capable of blocking the interaction between one or more IKKs such as IKKalpha or IKKbeta, and NEMO. The antiinflammatory compound further comprises at least one membrane translocation domain. The compounds are useful for treating inflammatory disorders, autoimmune diseases, osteoporosis, cancer, Alzheimer's disease, atherosclerosis, viral infections. Ataxia telangiectasia, and for transplantation detection. The compounds of the invention block NF-kappaB induction by IKK but do not inhibit the invention block NF-kappaB induction by IKK but do not inhibit
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human; antiinflammatory compound; NEMO binding domain; NBD; IKKbeta; IkappaB kinase-beta; IkappaB kinase-alpha; IKKalpha; NF-kappaB; nuclear factor-kappaB induction; inflammatory disorder;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human NEMO binding site (NBD) mutant peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Novel antiinflammatory peptide compounds comprising NEMO binding domain, useful for modulating NF-kappaB induction in a cell and i treating NF-kappaB-mediated inflammation disorders e.g., asthma, psoriasis, vasculitis -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Homo sapiens
Synthetic.
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22-AUG-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     02-MAY-2001; 2001US-0847940.
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atherosclerosis; viral infection; Ataxia telangiectasia;
                                                                                                                                                                                                                                                    Seguence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 2003-209142/20.
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                                                                                                                                                                                                                                                                                                                                         telangiectasia, and for transplantation detection.
the invention block NF-kappaB induction by IKK but
the basal activity of NF-kappaB. ABU08418-ABU08432
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (/MYAM/)
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GHOSH
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                                                         LDASAL 6
                                                                                                                                                                                                                                                    6 AA;
                                                                                                                                                                                                                                                                                                              peptides.
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2000US-0643260.
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                                                                                                                                                         100.0%;
                                                                                                                                                         Score 26; DB 24;
Pred. No. 9.3e+05;
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(UYYA) UNIV YALE.

02-MAY-2000; 2000US-201261P 22-AUG-2000; 2000US-0643260

02-MAY-2001; 2001WO-US40654

Misc-difference

note= "Wildtype Trp substituted by Ala"

WO200183547-A2

Misc-difference

Location/Qualifiers

'note= "Wildtype

dıl

substituted by Ala"

Synthetic

antiarthritic; osteopathic; antiulcer; mutant; mutein. dermatological; antibacterial; antipsoriatic; antirheumatic;

Modulating NF-kappaB induction in a cell, useful for treating e.g. inflammatory disorders, osteoporosis and cancer, comprises contacting cell with an anti-inflammatory compound comprising at least one NEMO binding domain

WPI; 2002-179350/23.

мау мJ,

Ghosh S;

Claim 23; Fig 5; 82pp; English

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CC The invention relates to modulating NF-kappaB (NF-kB) induction in a cell comprises contacting a cell with an anti-inflammatory compound CC (ABB08755-ABB08742) comprising at least one NEMO binding domain CC (ABB7313). The compound has acts through selective inhibition of CC with IKKbeta at the NEMO binding domain. Blockage of IKKbeta NEMO compound the results in inhibition of IKKbeta kinase activation and contested in results in inhibition of IKKbeta kinase activation and contested phosphorylation of IKKbeta kinase activation and activation of IKKbeta kinase activation of IKKbeta kinase activation of IKKbeta kinase activation of IKKbeta kinase activation and activation of IKKbeta kinase acti
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RESULT 5
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Antiinflammatory; antiasthmatic; cytostatic; antipsoriatic; nootropic; antiirheumatic; antiarthritic; osteopathic; antibacterial; virucide; immunosuppressive; dermatological; neuroprotective; antiatherosclerotic; antiallergic; membrane translocation domain; NEMO binding domain; eczema; cytokine; NFkappaB; IkappaB kinase beta; IKKbeta; cancer; psoriasis; rheumatoid arthritis; osteoarthritis; inflammatory bowel disease; autoimmune disorder; multiple sclerosis; transplant rejection; osteoporossis; Alzheimer's disease; atherosclerosis; viral infection; ataxia telangiectasia; allergy; anaphylaxis; arthritis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence
   AAM48628-AAM48645), comprising a membrane translocation d
(AAM48620-AM48627 or AAM48646-AAM48651) which comprises
amino acid residues, fused to a NEMO binding sequence
(AAM48525-AAM48619). The antiinflammatory compounds have
                                                                                                                                                                                                                                                                                             Novel antiinflammatory compound comprising membrane translocation domain fused to NEWO binding sequence, useful for blocking nuclear factor kappaB activation, and for treating asthma, lung inflammation,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        02-MAY-2000; 2000US-201261P.
22-AUG-2000; 2000US-0643260.
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                                                                                                                                                                                                Example 5; Fig
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                                                                                                                                relates to an antiinflammatory compound
                                                                                                                                                                                                5; 88pp; English.
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Pred. No.
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antiinflammatory compounds have antiasthmatic

Novel antiinflammatory peptide compounds comprising NEWO binding domain, useful for modulating NF-kappaB induction in a cell and treating NF-kappaB-mediated inflammation disorders e.g., asthma, psoriasis, vasculitis -

for

22; Fig 5A; 47pp;

English

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nuclear factor-kappaB induction; inflammatory disorder;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence
                                                                                                                                                                                               02-MAY-2000;
22-AUG-2000;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human mutant NEMO binding site (NBD) peptide.
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                                                                                                  WPI; 2003-209142/20.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                          NF-kappaB;
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RESULT 7
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The present invention relates to antiinflammatory compounds comprising NEMO binding domain (NED) peptides. The NEMO binding domains are found on IkappaB kinase-beta (IKKbeta) and IkappaB kinase-alpha (IKKalpha) proteins. The antiinflammatory compounds of the invention are useful for modulating nuclear factor-kappaB (NF-kappaB) induction in a cell, where the compounds are capable of blocking the interaction between one or more IKKs such as IKKalpha or IKKbeta, and NEMO. The antiinflammatory compound further comprises at least one membrane translocation domain. The compounds are useful for treating inflammatory disorders, autoimmune disease, osteoporosis, cancer, Alzheimer's disease, atherosclerosis, viral infections, Ataxia telanglectasia, and for transplantation detection. The compounds of the invention block NF-kappaB induction by IKK but do not inhibit the basal activity of NF-kappaB. The present sequence represents
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             proteins from various Mycobacterium species microorganisms. The encoding nucleotide sequences can be used as primers and probes for methods for detecting and identifying mycobacteria, especially belonging to the M. tuberculosis complex. The encoded proteins can be used in vaccines for immunisation against a bacterial or viral infection.
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N-PSDB; AAX34203.
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                                                                                                             Sequences AAY04742-Y05000 and AAY07201-Y07204 represent secreted
                                                                                                                                               Claim 32; Fig 41D; 309pp; French
                                                                                                                                                                                                                                                                                                                    Guigueno
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14-AUG-1997;
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ation; detection; vaccine; immunisation; infection.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mycobacterial DNA vectors containing reporter constructs identifying coding or promoter sequences involved in infection-associated protein expression
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N-PSDB; AAX34200.
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14-AUG-1997;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim 32;
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N-PSDB; AAX34204.
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Guigueno A;
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14-AUG-1997;
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  25-FEB-1999;
                    25-FEB-2000; 2000EP-0301439.
                                        06-SEP-2000
                                                            EP1033405-A2
                                                                               Arabidopsis thaliana
                                                                                                          Protein identification; signal transduction pathway; metabolic pathway; hybridiaation assay; genetic mapping; gene expression control; promoter;
                                                                                                                                        Arabidopsis thaliana protein fragment SEQ ID NO: 43425.
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Similarity 100.0%;
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25-FEB-1999;
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N-PSDB; AAI81838.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Claim 20; SEQ ID NO 15799; 1399pp +
                                                                                                                                                                                                                                                                                                                                                                                                                       protein identification; signal transduction pathway; metabolic pathway;
hybridisation assay; genetic mapping; gene expression control; promoter;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Arabidopsis thaliana protein fragment SEQ ID NO: 43424.
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RESULT 14
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26-0CT 1999;
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07-APR-2000;
03-AUG-2000;
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Tateishi
The present invention provides a number of nucleotide and protein sequences from the Coryneform bacterium Corynebacterium glutamicum. These are useful for identifying the mutation point of a gene derived from a mutant of coryneform bacterium, measuring expression amount and analysing the expression profile or expression pattern of a gene derived from coryneform bacterium, and identifying a homologue of a gene derived from coryneform bacterium. Coryneform bacteria are useful for producing amino acids, nucleic acids, vitamins, saccharides and organic acids, particularly L-lysine. The present sequence is a protein described in the exemplification of the invention.

Note: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from the
                                                                                                                                                                                                                                                                       Novel polymucleotides derived from Coryneform bacteria, for identifying mutation point of a gene, measuring expression of a gene, analysing expression profile or pattern of a gene and identifying homologous gene -
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                                                                                                                                                                                                                             Claim 17; SEQ ID NO: 4338; 246pp + Sequence Listing; English
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Senoh A,
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2000JP-0280988.
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99US-0161992.
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99US-0162142.
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Pred. No. 1.4e+02;
; Mismatches 0;
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Ozaki A;
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21-MAY-1999;
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28-MAY-1999;
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14-MAY-1999;
14-MAY-1999;
18-MAY-1999;
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28-APR-1999;
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04-MAY-1999
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29-MAR-1999;
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09-MAR-1999;
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                                                                                                                                                                                                                                                                    11-MAY-1999,
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                                                                                                                                                             19-MAY-1999;
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                                     99US-0123180

99US-0125788

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99US-0130449

99US-0132048

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SXS
Arabidopsis thaliana.
                                        Protein identification; signal transduction pathway; metabolic pathway; hybridisation assay; genetic mapping; gene expression control; promoter;
                                                                                     Arabidopsis thaliana protein fragment SEQ ID
                                                                                                                                                                                                                                                                                                                                                                                              Patent Office
                             sequence
                                                                                                                                                                                                                                                                                                          Conservative
                                                                                                                  (first
                                                                                                                                                                             Protein; 271 AA
                                                                                                                  entry)
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 99US-0137724.
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99US-014536.
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Best Local Similarity 100.0%;
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Pred. No. 2.4e+02;
Mismatches 0;
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Search completed: February 18, 2004, 14:26:18 Job time: 23.7763 secs